

Fig. 1

	10	20	30	40	50	60
ulbp2	MAAAAATKIL	LCLPLLLLLS	GWSRAGRADP	HSLCYDITVI	PKFRPGPRWC	AVQGQVDEKT

r15	MAAAASPAFL	LRLPLLLLLS	SWCRTGLADP	HSLCYDITVI	PKFRPGPRWC	AVQGQVDEKT
	10	20	30	40	50	60
	70	80	90	100	110	120
ulbp2	FLHYDCGNKT	VTPVSPLGKK	LNVTTAWKAQ	NPVLREVVDI	LTEQLRDIQL	ENYTPKEPLT

r15	FLHYDCGSKT	VTPVSPLGKK	LNVTTAWKAQ	NPVLREVVDI	LTEQLLDIQL	ENYIPKEPLT
	70	80	90	100	110	120
	130	140	150	160	170	180
ulbp2	LQARMSCEQK	AEGHSSGSWQ	FSFDGQIFLL	FDSEKRMWTT	VHPGARKMKE	KWENDKVVAM

r15	LQARMSCEQK	AEGHSGGSWQ	LSFDGQIFLL	FDSENRMWTT	VHPGARKMKE	KWENDKDMTM
	130	140	150	160	170	180
	190	200	210	220	230	240
ulbp2	SFHYFSMGDC	IGWLEDFLMG	MDSTLEPSAG	APLAMSSGTT	QLRATATTLI	LCCLLIILPC

r15	SFHYISMGDC	TGWLEDFLMG	MDSTLEPSAG	-----GTV		
	190	200	210	220	230	

ulbp2 FILPGI

r15

Fig. 2A

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U1bp2: 1  atggcagcagccgcccgtaccaagatccttctgtgcctcccgttctgctcctgtgtcc 60
          ||||| |||||  ||  ||||| ||||| ||||| ||||| |||||
rl5: 1  atggcagcagccgcccagccccggttccttctacgcctcccgttctgctcctgtgtcc 60

ulbp2: 61  ggctgggtccgggctgggcgagccgacctcactctcttctgtatgacatcacgcgtcatc 120
          ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
rl5: 61  agctgggtgcaggaccgggctggccgacctcactctcttctgtatgacatcacgcgtcatc 120

ulbp2: 121 cctaagttcagacctggaccacgggtggtgtgcggttcaaggccaggtggatgaaaagact 180
          ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
RL5: 121 cctaagttcagacctggaccacgggtggtgtgcggttcaaggccaggtggatgaaaagact 180

U1bp2: 181 tttcttcactatgactgtggcaacaagacagtcacacctgtcagtcacctggggaagaaa 240
          ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
RL5: 181 tttcttcactatgactgtggcagcaagacagtcacacctgtcagtcacctggggaagaaa 240

U1bp2: 241 ctaaatgtcacaacggcctggaaagcacagaacccagtactgagagaggtggtggacata 300
          ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
RL5: 241 ctaaatgtcacaacggcctggaaagcacagaacccagtactgagagaggtggtggacata 300

U1bp2: 301 cttacagagcaactgcgtgacattcagctggagaattacacaccaaggaacccctcacc 360
          ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
RL5: 301 cttacagagcaactgcttgacattcagctggagaattacataaccaaggaacccctcacc 360

U1bp2: 361 ctgcaggcaaggatgtcttgtgagcagaaagctgaaggacacagcagtggtcttggcag 420
          ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
RL5: 361 ctgcaggccaggatgtcttgtgagcagaaagccgaaggacacggcagtggtcttggcag 420

U1bp2: 421 ttcagtttcgatgggcagatcttctcctcttcttctgactcagagaagagaatgtggacaacg 480
          ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
RL5: 421 ttcagtttcgatggacagatcttctcctcttcttctgactcagaaaacagaatgtggacaacg 480

U1bp2: 481 gttcatcctggagccagaaagatgaaagaaaagtgggagaatgacaagggttggccatg 540
          ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
RL5: 481 gttcatcctggagccagaaagatgaaagaaaagtgggagaatgacaaggatgacatg 540

U1bp2: 541 tccttccattactttctcaatgggagactgtataggatggcttgaggacttcttggatgggc 600
          ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
RL5: 541 tccttccattacatctcaatgggagactgcacaggatggcttgaggacttcttggatgggc 600

U1bp2: 601 atggacagcaccctggagccaagtgcaggagcaccactcgccatgtcctcaggcacaacc 660
          ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
RL5: 601 atggacagcaccctggagccaagtgcaggaggcacagtctga-----

U1bp2: 661 caactcagggccacagccaccaccctcatccttctgtgcctcctcatcatcctccctgc 720
          ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
RL5: -----

U1bp2: 721 ttcattcctccctggcatctga
          ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
RL5: -----

```

Fig. 2B

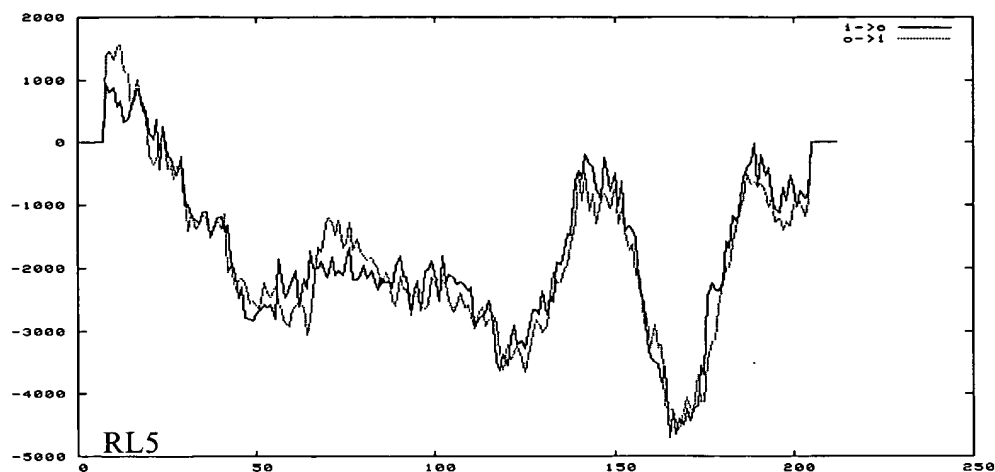


Fig. 3A

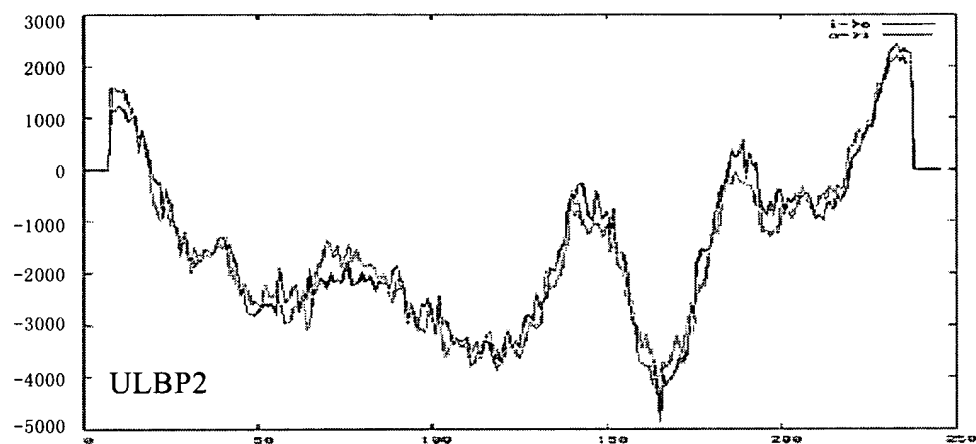


Fig. 3B

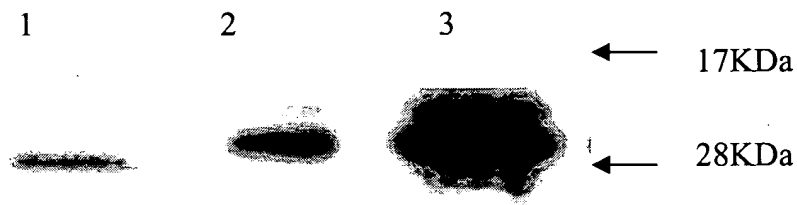


Fig. 4

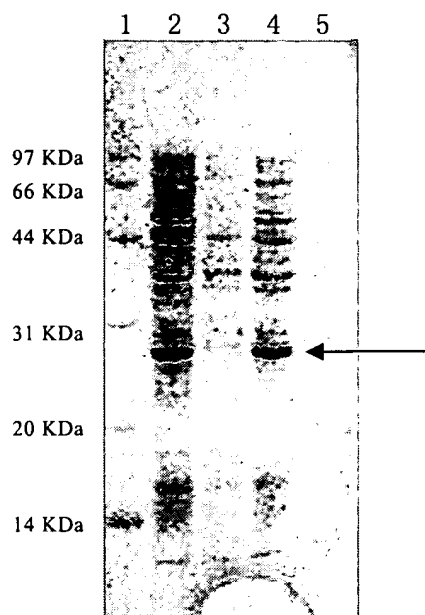


Fig. 5

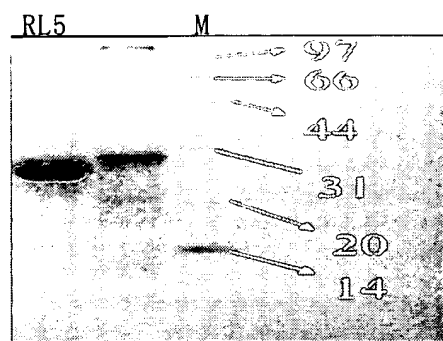
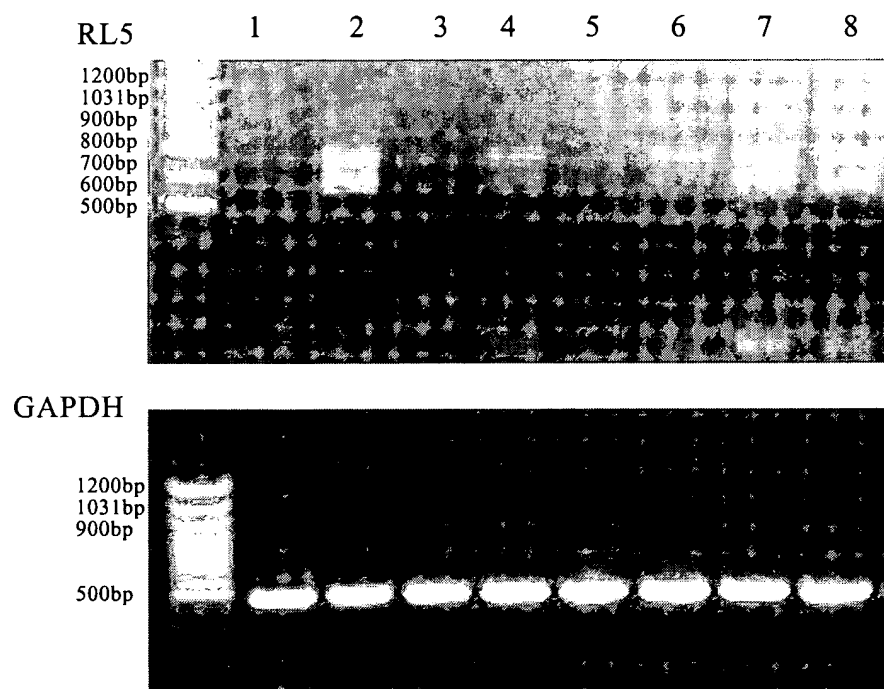


Fig. 6



1=stomach; 2=stomach cancer; 3=colon; 4=colon cancer
5=breast; 6=breast cancer; 7=kidney; 8=kidney cancer

Fig. 7

RT-PCR

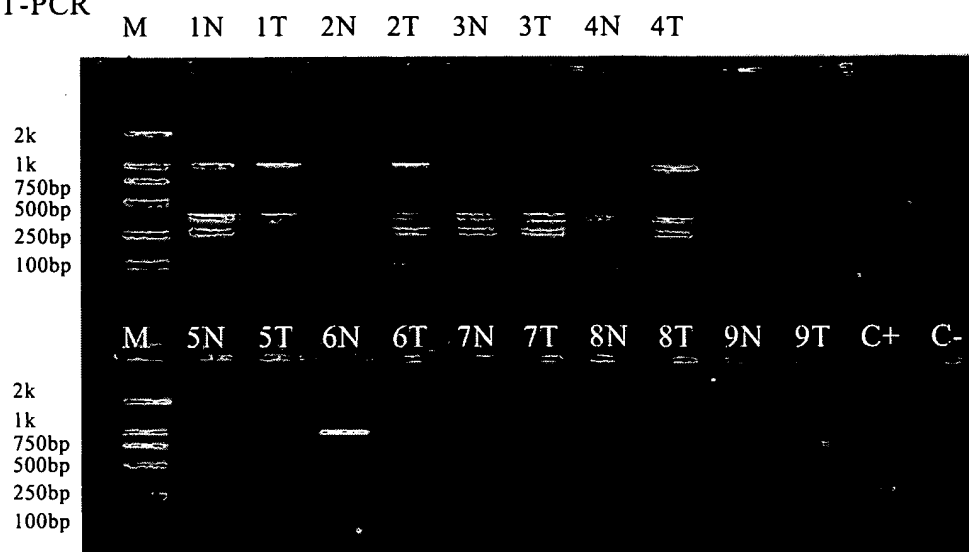


Fig. 8A

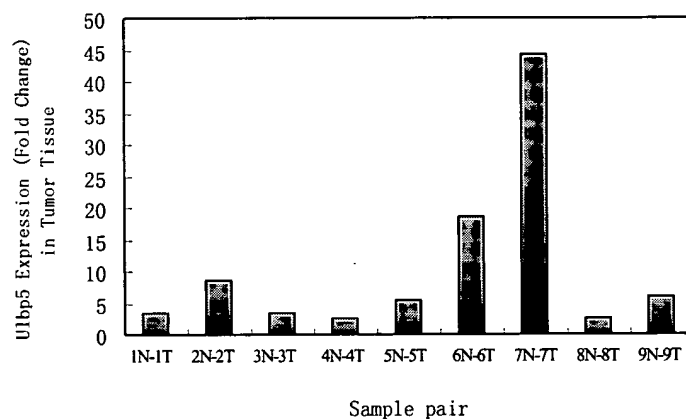


Fig. 8B

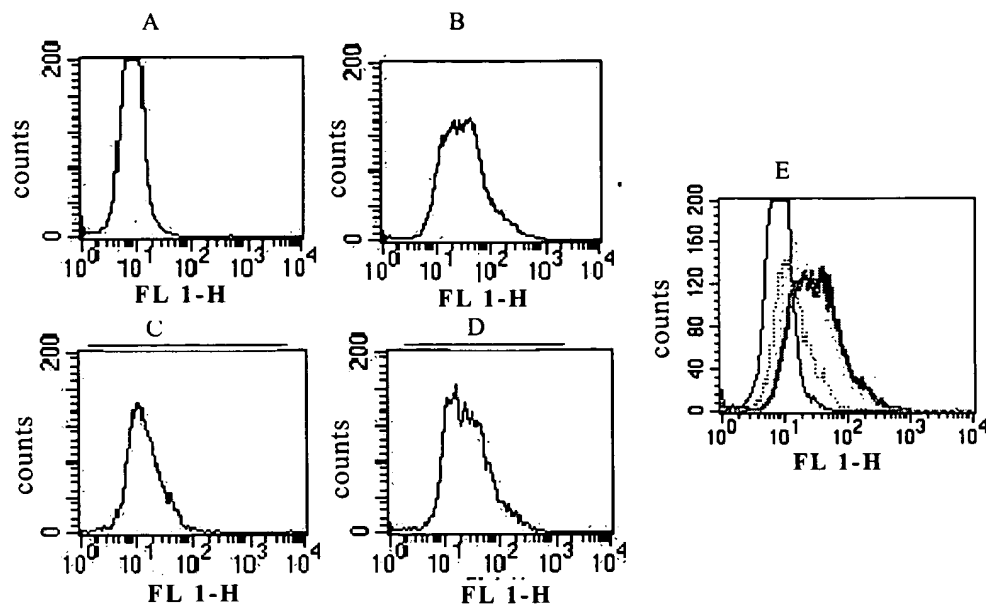


Fig. 9